

FIG. 1

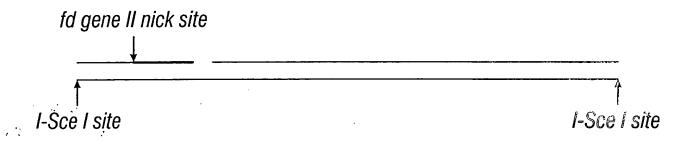
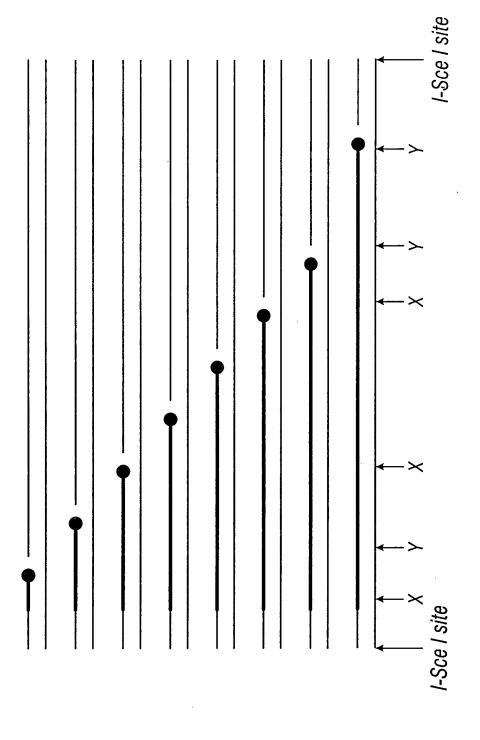


FIG. 2

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F16. 3

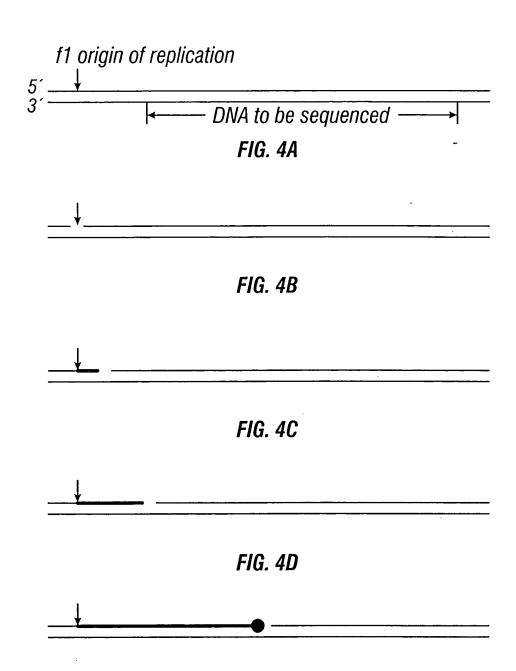


FIG. 4E

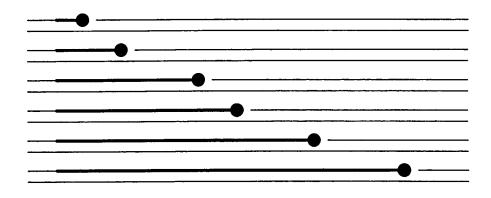


FIG. 4F

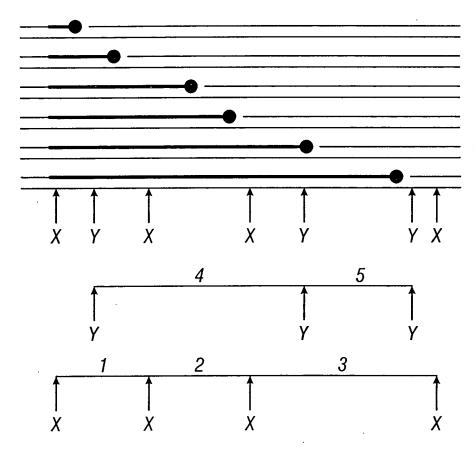


FIG. 4G

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FIG. 4H

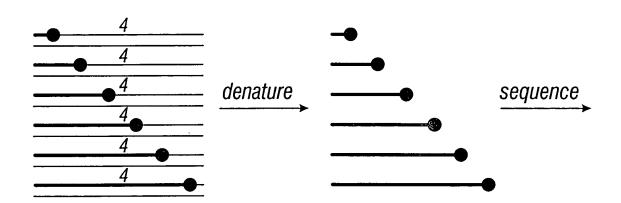


FIG. 41

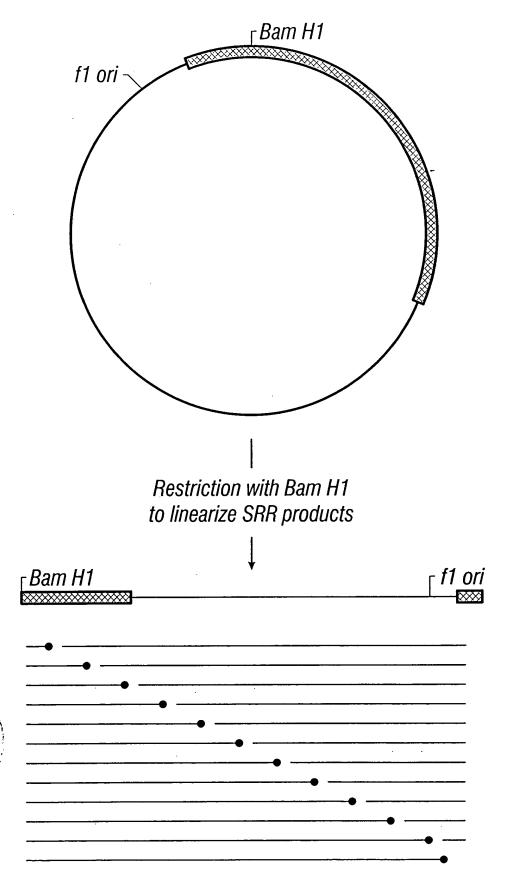
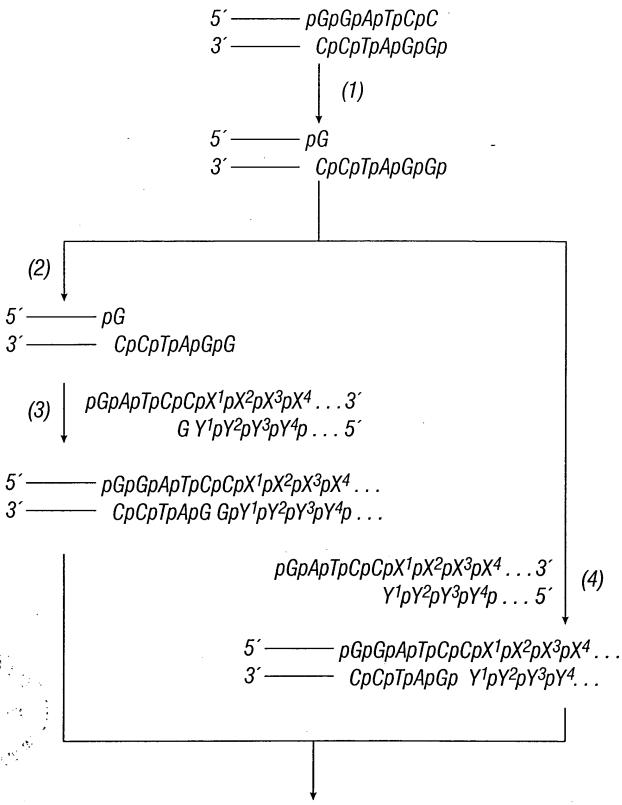


FIG. 5



STRAND REPLACEMENT REACTION

FIG. 6

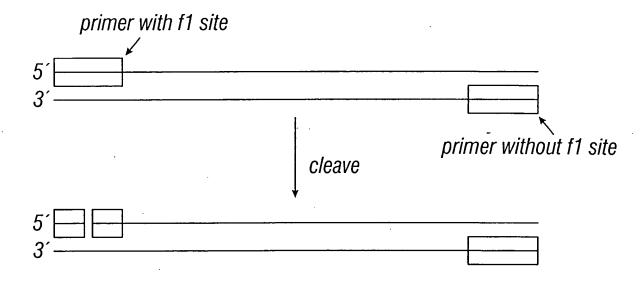


FIG. 7A

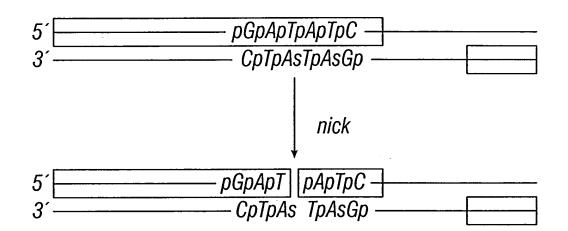
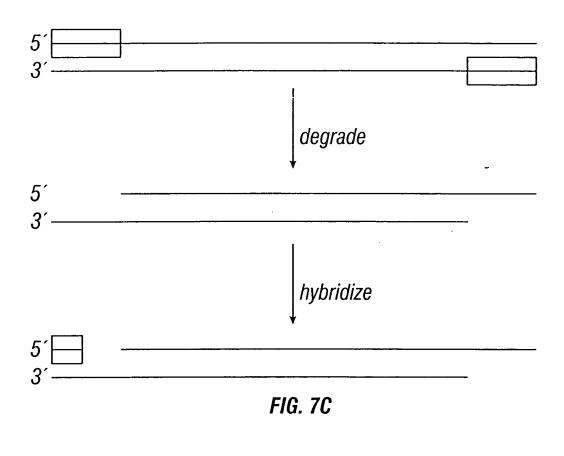
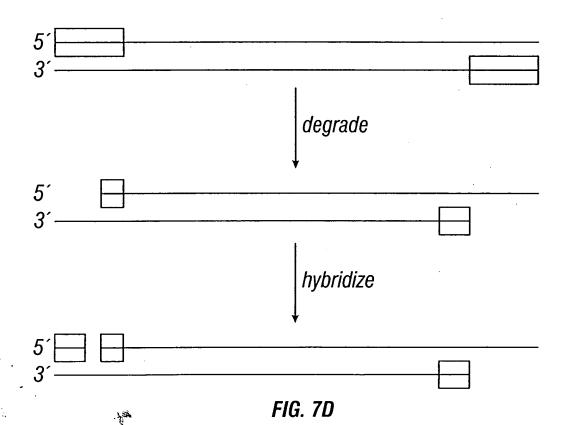


FIG. 7B





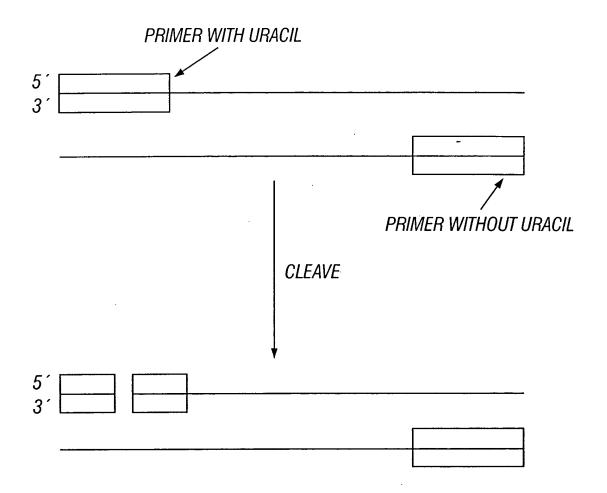


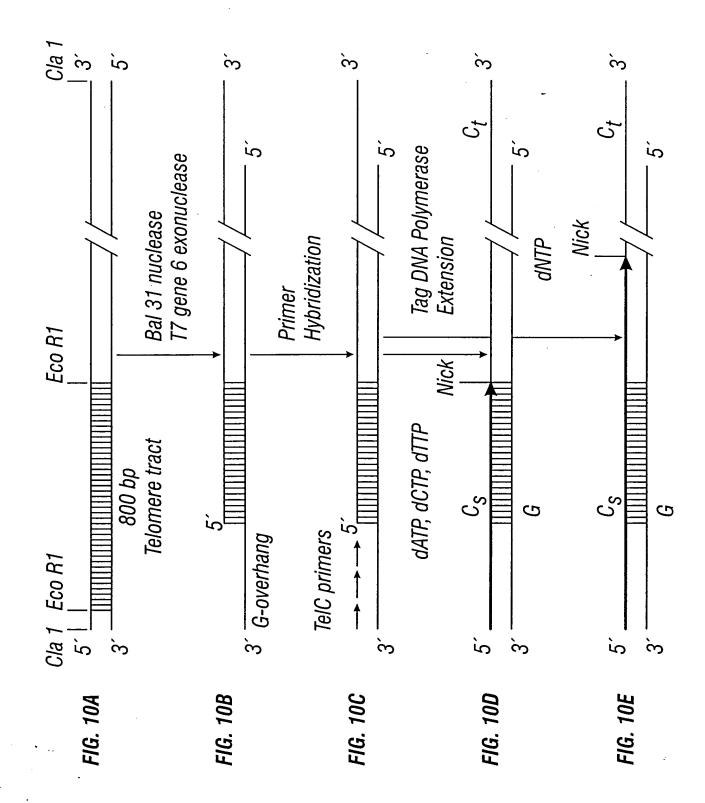
FIG. 7E

₽Ú.	detectable features to be mapped
template DNA	DNA
SR reacti	SR reaction time (min)
with dUTF	with dUTP with dTTP
0	
10	1
20	1
30	1
40	1
>	= thymidine-containing DNA synthesized by SR = uridine-containing DNA synthesized by SR

FIG. 8

nick site ↓					= DNA with labeled dTTP to be used for array hybridization
fd nick site ↓					

FIG. 9



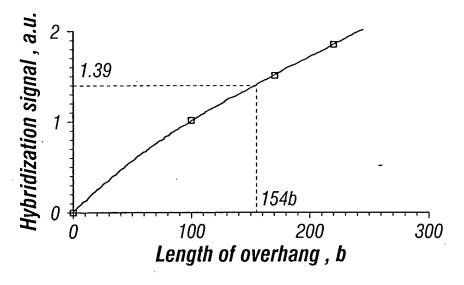
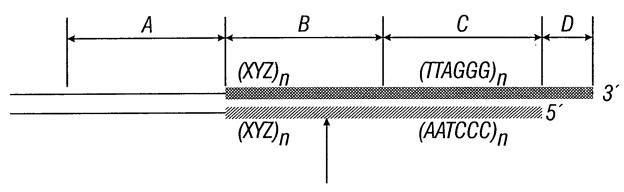
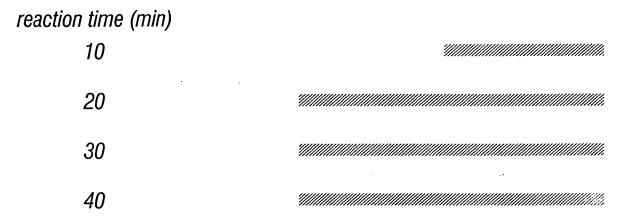


FIG. 11



site of first guanine in the C-rich strand



=DNA synthesized by SR using only dATP, dTTP, and dCTP

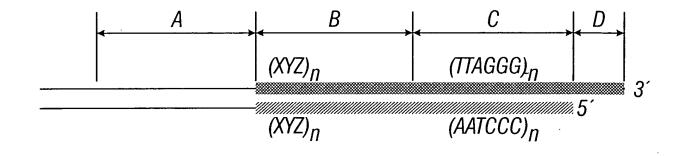


FIG. 13

FIG. 14B

B 123456789 123456789

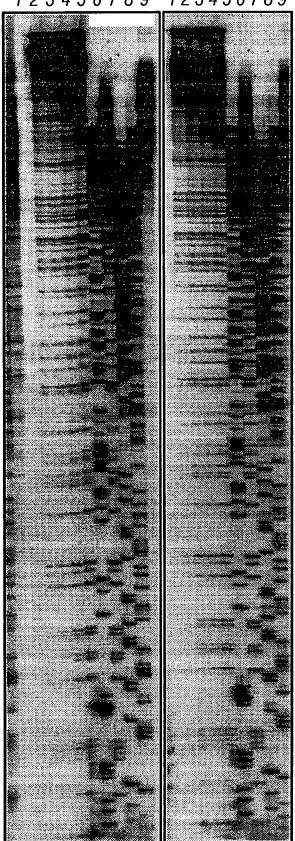


FIG. 14A

Complyto copos

OF SHOWN)	15A 15B FIG. 15	
PCR-AMPLIFIED WITH DETECTION TAG AT 5' END C PRIMER X. NUMBERS LABEL THE 12 UNKNOWN BASES. RANDOM DEGRADATION (ONLY DAMAGED UPPER STRAND S	O	EXPOSE 3'OH AT DAMAGE SITES O
\$'OA-C-T-A-C-T-G-G-T-T-T-A3' 3'A-T-G-A-T-A-C-C-A-A-T-T-T5' 1 2 3 4 5 6 7 8 9101112 PRIMER X UNKNOWN DNA SEQUENCE PRIMER Y	O	O

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INCORPORATE BIOTINYLATED DDTTP AT POSITIONS OPPOSITE ADENINE IN TEMPLATE STRAND	OT-A-C-T-A-T G-T-T-A-C-C-A-A-C-C-A-A-A-T-C-C-A-A-A-T-A-C-C-A-A-T-A-C-C-A-A-A-T-A-C-C-A-A-T-A-T	IMMOBILIZE BIOTINYLATED STRANDS AND REMOVE NON-BIOTINYLATED STRANDS OT-A-C-T-A-T-G-G-T. OT-A-C-T-A-T-G-G-T-T.	RELEASE BIOTINYLATED STRANDS, SEPARATE BY ELECTROPHORESIS, AND DETECT TAGGED PRIMERS (DARK BARS REPRESENT POSITIONS OF THYMINE) 7 8 9 10 11 12 T T T T
		T. -A-C-T. -T-A-T.	7. P. F.
	T-T-A T-T-A-T A-A-T T-T-A-T T-T-A-T T-T-A-T T-T-A-T T-T-A-T T-T-A-T A-A-T A-A-T	O	4 2222 F1
		0	m
	4	O	1 2
· .:	T.A-C-TA-T-G-AT-AT-AT-AT-A-C-TA-T-G-AT-A-C-TA-T-G-AT-A-C-TA-T-G-A		
• •			
			•

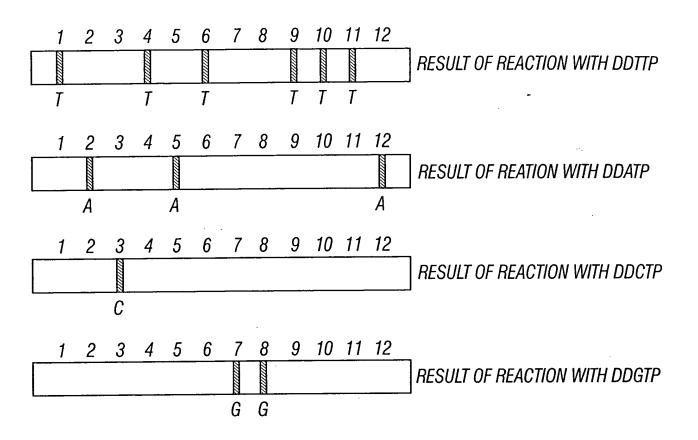


FIG. 16A

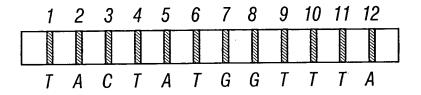


FIG. 16B

PCR-AMPLIFIED WITH DETECTION TAG AT 5' END OF PRIMER X. NUMBERS LABEL THE 12 UNKNOWN BASES. RANDOM DEGRADATION (ONLY DAMAGED UPPER STRAND SHOWN)	O	EXPOSE 3'OH AT DAMAGE SITES OT-A-C-T-A-T G-T-T-A-C-C-O-O-O-O-O-O-O-T-A-C-T-A-T-G T-T-T-A-C-C-O-O-O-O-O-O-O-O-O-O-O-O-O-O-O-O-O
5'OT-A-C-T-A-T-G-G-T-T-T-A3' 3'A-T-G-A-T-A-C-C-A-A-T5' 1 2 3 4 5 6 7 8 9101112 PRIMER X UNKNOWN DNA SEQUENCE PRIMER Y	O	O

DOMOKULO OMOTOK

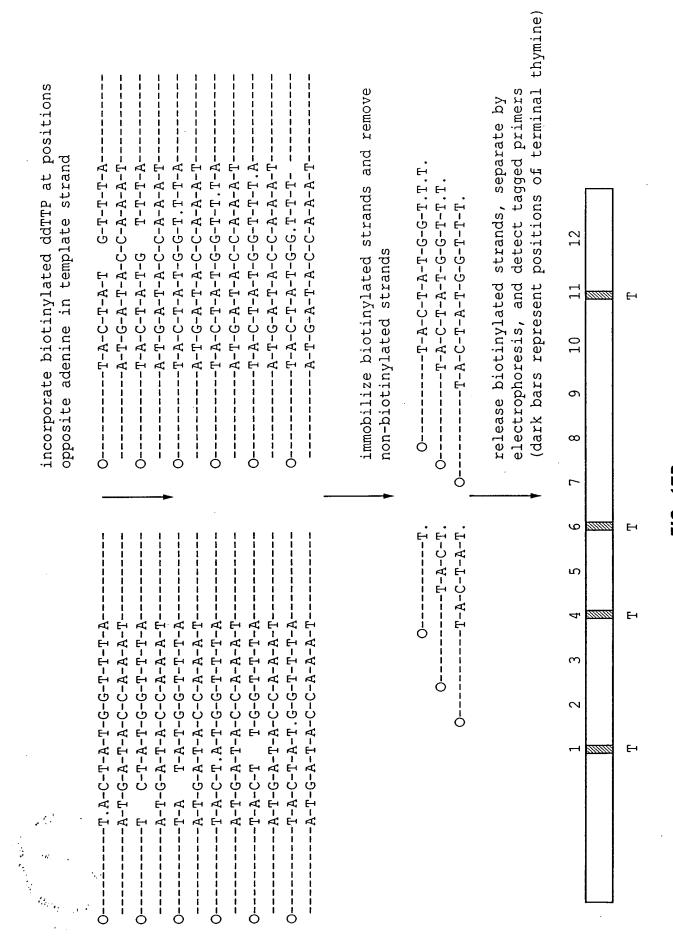


FIG. 17B

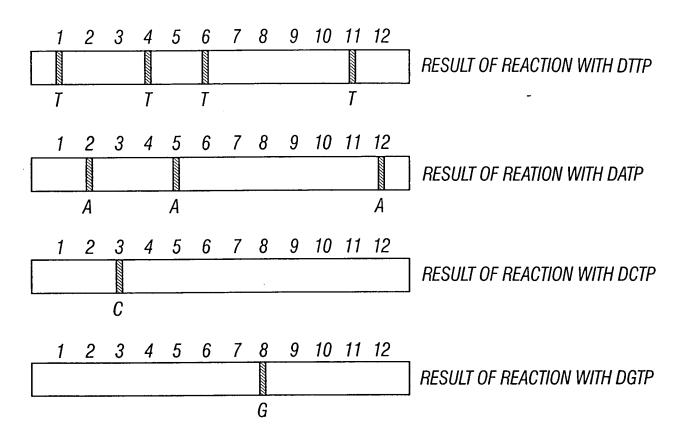


FIG. 18A

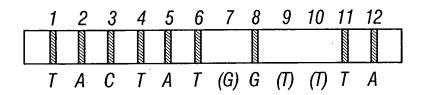


FIG. 18B

FIG. 19

INCORPORATE TAGGED DDTTP AT POSITIONS

OPPOSITE ADENINE IN TEMPLATE STRAND

FIG. 19B

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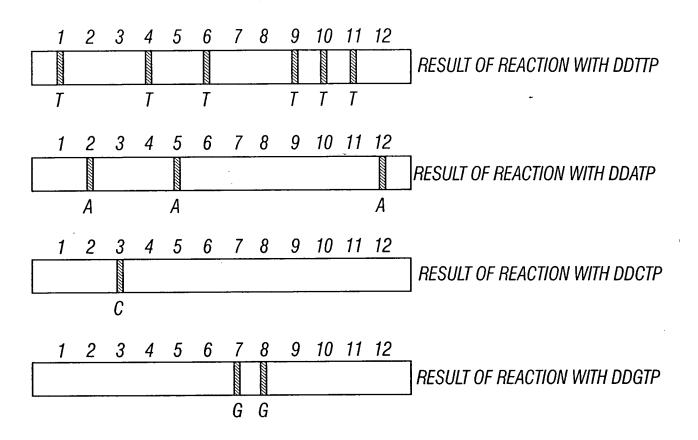


FIG. 20A

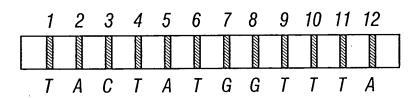


FIG. 20B

ilize, and expose OH in Fig. 5.	O	Block ends opposite T, G' & C w (shown in bold letters), remove OT-A-C-T-A-T-G-G-T-T-O
mmobi s as	O	O

-A-T----

- A - T - A -

with ddATP, ddGTP, ddCTP e ddNTPs, then add dTTP. 21A 21B O------T-A-C-T-A-T-G-G-T-T-T-A----------A-T-G-A-T-A-C-C-A-A-T----1-A-T-----O------T-A-C-T-A-T-G-G-T-T-T-A--------T-A----------------------------A-T-G-A-T-A-C-C-A-A-T------------------T-A-----T-A---

FIG. 21A

O------T-A-C-T-A-T-G-G-T-T-T-A-

----A-T-G-A-T-A-C-C-A-A-T---

-----A-T-G-A-T-A-C-C-A-A-A-T-----

FIG. 21B

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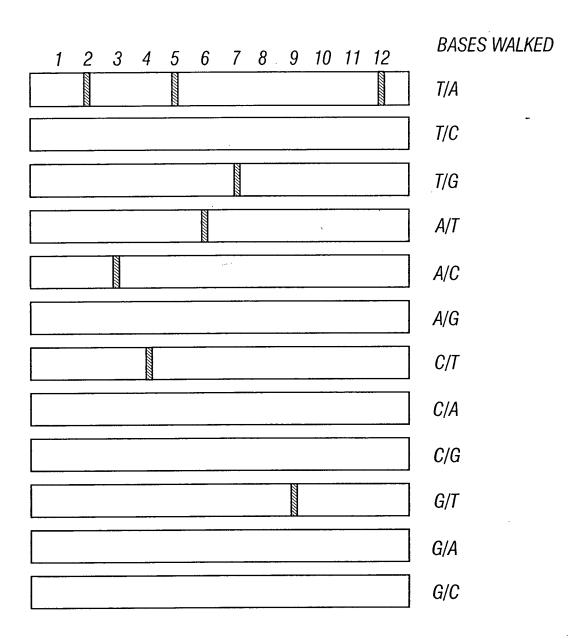


FIG. 22A

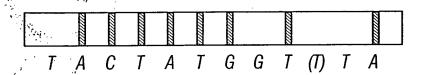


FIG. 22B

at random sites as in Fig.

НО <u>-</u>

FIG. 23

23A

23B

H

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EH

romorum.cmc.

Block ends opposite A, G & C with ddTTP, ddGTP, ddCTP (shown in bold), remove ddNTPs, then add dATP.	OA-T-A-T-G-G-T-T-T-A-T-A	G-A-T-A-C-C- C-T-A-T-G-G- G-A-T-A-C-C-C-C-C-C-C-C-C-C-C-C-C-C-C-C-C		A-T-G-A-T-A-C-C-A-A-A-T Block ends opposite A, G & C with ddTTP, ddGTP, ddCTP Block ends opposite T, G & C with ddATP, ddGTP, ddCTI (shown in bold), remove ddNTPs, then add tagged ddTTI	 O	, size-separate, and detect strands with tagged terminal T $$7\ 8\ 9\ 10\ 11\ 12$
0	OT-A-C-T-A-T-G-G-T-T-T-A-T-A	A-T-G-A-T-A-C-C-A-A-A- T-A-C-T-A-T-G-G-T-T-T- A-T-G-A-T-A-C-C-A-A-A-	-A-C-I-A-I-G-G-I-I-I- -T-G-A-T-A-C-C-A-A-A- -A-C-T-A-T-G-G-T-T-T- -T-G-A-T-A-C-C-A-A-A- -A-C-T-A-T-G-G-T-T-T-	A-T-G-A-T-A-C-C-A-A-T	 O	Remove all non-immobilized DNA, then release, 1 2 3 4 5 6

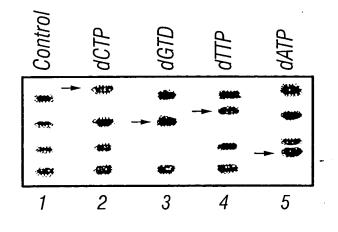
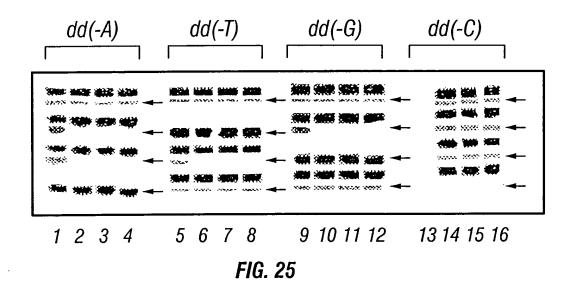


FIG. 24



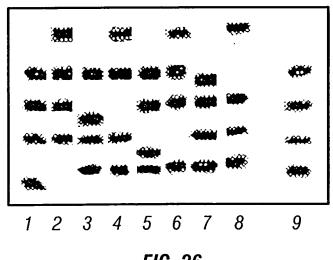


FIG. 26

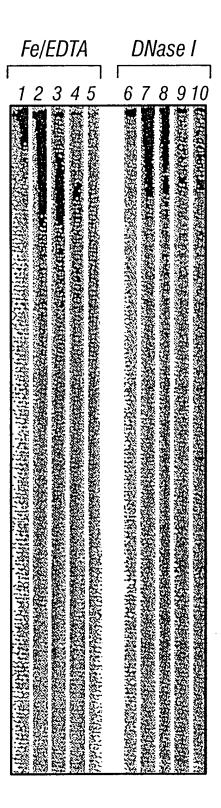
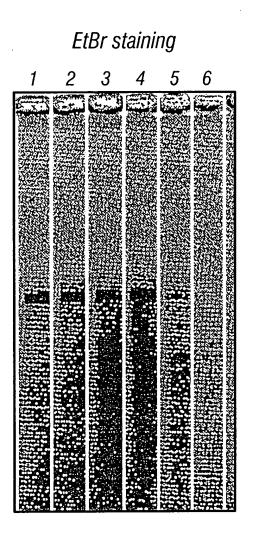


FIG. 27



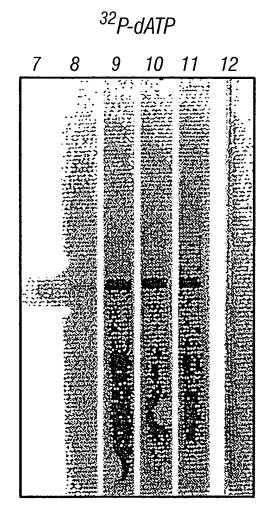


FIG. 28A

FIG. 28B

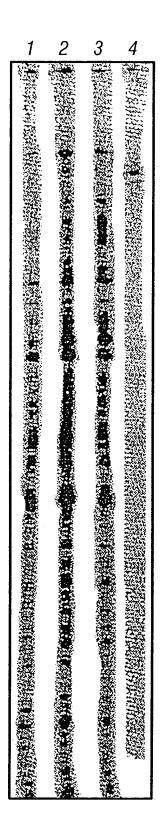
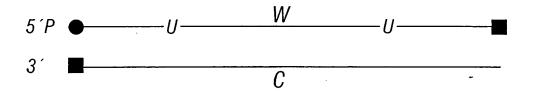


FIG. 29



- -5´-PHOSPHATE
- - 3´ DIDEOXYNUCLEOTIDE OR NH₃ GROUP

FIG. 30A

5 '		X	3´ OH 16 C-XY OLIGOS
	X, Y AND Z ARE A, T, G OR C		

FIG. 30B

1			ļ	ļ	
				OM DOUBLE K INCORPOR	
			XYZ		
	XYZ	XYZ			
XYZ					
		MULTI-E	BASE SEL	ECTION.	
XYZ	XYZ	XYZ	XYZ		
	SIZE SE	V PARATION			

FIG. 31

A STATE OF THE STA

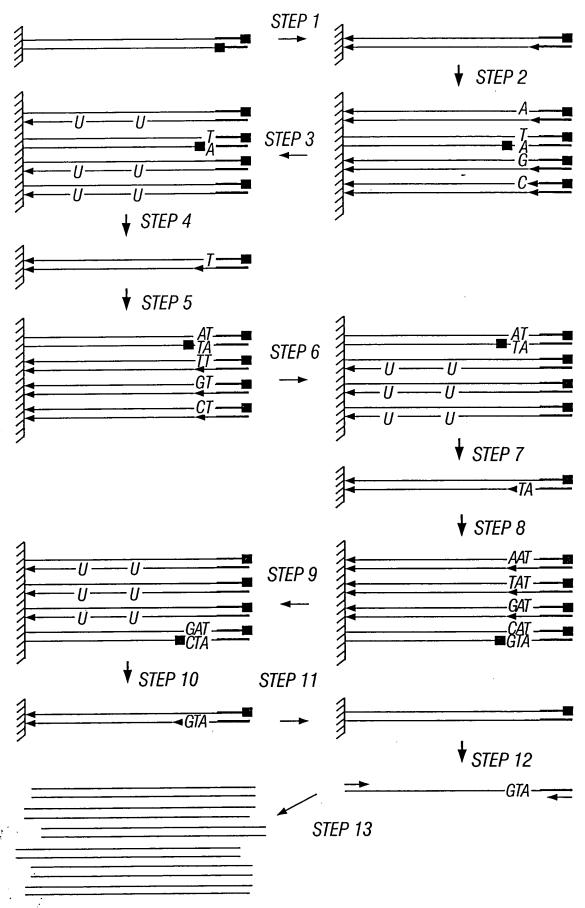


FIG. 32

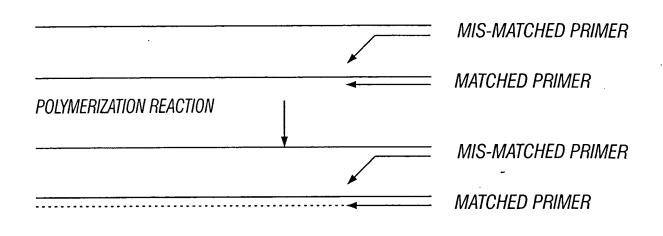


FIG. 33A

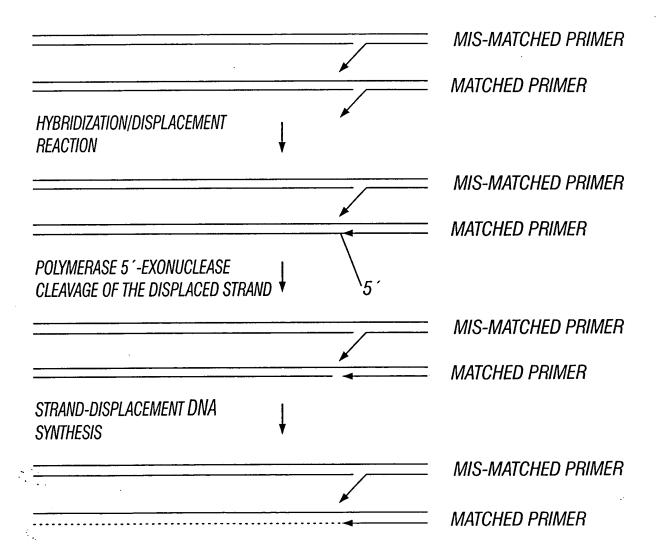


FIG. 33B